OIPE

RAW SEQUENCE LISTING   DATE: 01/30/2002
PATENT APPLICATION: US/10/044,807   TIME: 15:44:46

Input Set : A:\LEX-0298-USA SEQLIST.txt
Output Set: N:\CRF3\01302002\J044807.raw

ENTERED

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   4 <110> APPLICANT: Yu, Xuanchuan
   5        Turner, C. Alexander Jr.
   7 <120> TITLE OF INVENTION: Novel Human Protease and Polynucleotides Encoding the Same
   9 <130> FILE REFERENCE: LEX-0298-USA
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/044,807
C--> 11 <141> CURRENT FILING DATE: 2002-01-11
  11 <150> PRIOR APPLICATION NUMBER: US 60/261,684
  12 <151> PRIOR FILING DATE: 2001-01-12
  14 <160> NUMBER OF SEQ ID NOS: 2
  16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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  20 <212> TYPE: DNA
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  26 ccatggagtg aatgctcacg cacctgcggg ggtggggcct cctactctct gaggcgctgc    180
  27 ctgagcagca agagctgtga aggaagaaat atccgataca gaacatgcag taatgtggac    240
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  29 caccatggcc agttttatga atggcttcct gtgtctaatg accctgacaa cccatgttca    360
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  35 ggtcctgatc acttatatct ggaaaccaaa accctccagg ggactaaagg tgaaaacagt    720
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  37 gacaaagaga tactgagaat ggctggacca ctcacagcag atttcattgt caagattcgt    840
  38 aactcgggct ccgctgacag tacagtccag ttcatcttct atcaacccat catccaccga    900
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  40 tcggctgagt gctacgatct gaggagcaac cgtgtggttg ctgaccaata ctgtcactat   1020
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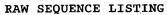
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126 Asp Gly Leu Trp Asp Ala Trp Gly Pro Trp Ser Glu Cys Ser Arg Thr
127         35                  40                  45
128 Cys Gly Gly Gly Ala Ser Tyr Ser Leu Arg Arg Cys Leu Ser Ser Lys
129     50                  55                  60
130 Ser Cys Glu Gly Arg Asn Ile Arg Tyr Arg Thr Cys Ser Asn Val Asp
131 65                  70                  75                  80
132 Cys Pro Pro Glu Ala Gly Asp Phe Arg Ala Gln Gln Cys Ser Ala His
133                 85                  90                  95
134 Asn Asp Val Lys His His Gly Gln Phe Tyr Glu Trp Leu Pro Val Ser
135             100                 105                 110
136 Asn Asp Pro Asp Asn Pro Cys Ser Leu Lys Cys Gln Ala Lys Gly Thr
137         115                 120                 125
138 Thr Leu Val Val Glu Leu Ala Pro Lys Val Leu Asp Gly Thr Arg Cys
139     130                 135                 140
140 Tyr Thr Glu Ser Leu Asp Met Cys Ile Ser Gly Leu Cys Gln Ile Val
141 145                 150                 155                 160
142 Gly Cys Asp His Gln Leu Gly Ser Thr Val Lys Glu Asp Asn Cys Gly
143                 165                 170                 175
144 Val Cys Asn Gly Asp Gly Ser Thr Cys Arg Leu Val Arg Gly Gln Tyr
145             180                 185                 190
146 Lys Ser Gln Leu Ser Ala Thr Lys Ser Asp Asp Thr Val Val Ala Ile
147         195                 200                 205
148 Pro Tyr Gly Ser Arg His Ile Arg Leu Val Leu Lys Gly Pro Asp His
149     210                 215                 220
150 Leu Tyr Leu Glu Thr Lys Thr Leu Gln Gly Thr Lys Gly Glu Asn Ser
151 225                 230                 235                 240
152 Leu Ser Ser Thr Gly Thr Phe Leu Val Asp Asn Ser Ser Val Asp Phe
153                 245                 250                 255
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157          275               280               285
158 Val Gln Phe Ile Phe Tyr Gln Pro Ile Ile His Arg Trp Arg Glu Thr
159     290               295               300
160 Asp Phe Phe Pro Cys Ser Ala Thr Cys Gly Gly Gly Tyr Gln Leu Thr
161 305               310               315               320
162 Ser Ala Glu Cys Tyr Asp Leu Arg Ser Asn Arg Val Val Ala Asp Gln
163               325               330               335
164 Tyr Cys His Tyr Tyr Pro Glu Asn Ile Lys Pro Lys Pro Lys Leu Gln
165          340               345               350
166 Glu Cys Asn Leu Asp Pro Cys Pro Ala Ser Asp Gly Tyr Lys Gln Ile
167          355               360               365
168 Met Pro Tyr Asp Leu Tyr His Pro Leu Pro Arg Trp Glu Ala Thr Pro
169     370               375               380
170 Trp Thr Ala Cys Ser Ser Ser Cys Gly Gly Gly Ile Gln Ser Arg Ala
171 385               390               395               400
172 Val Ser Cys Val Glu Glu Asp Ile Gln Gly His Val Thr Ser Val Glu
173               405               410               415
174 Glu Trp Lys Cys Met Tyr Thr Pro Lys Met Pro Ile Ala Gln Pro Cys
175          420               425               430
176 Asn Ile Phe Asp Cys Pro Lys Trp Leu Ala Gln Glu Trp Ser Pro Cys
177          435               440               445
178 Thr Val Thr Cys Gly Gln Gly Leu Arg Tyr Arg Val Val Leu Cys Ile
179     450               455               460
180 Asp His Arg Gly Met His Thr Gly Gly Cys Ser Pro Lys Thr Lys Pro
181 465               470               475               480
182 His Ile Lys Glu Glu Cys Ile Val Pro Thr Pro Cys Tyr Lys Pro Lys
183               485               490               495
184 Glu Lys Leu Pro Val Glu Ala Lys Leu Pro Trp Phe Lys Gln Ala Gln
185          500               505               510
186 Glu Leu Glu Glu Gly Ala Ala Val Ser Glu Glu Pro Ser Phe Ile Pro
187          515               520               525
188 Glu Ala Trp Ser Ala Cys Thr Val Thr Cys Gly Val Gly Thr Gln Val
189     530               535               540
190 Arg Ile Val Arg Cys Gln Val Leu Leu Ser Phe Ser Gln Ser Val Ala
191 545               550               555               560
192 Asp Leu Pro Ile Asp Glu Cys Glu Gly Pro Lys Pro Ala Ser Gln Arg
193               565               570               575
194 Ala Cys Tyr Ala Gly Pro Cys Ser Gly Glu Ile Pro Glu Phe Asn Pro
195          580               585               590
196 Asp Glu Thr Asp Gly Leu Phe Gly Gly Leu Gln Asp Phe Asp Glu Leu
197          595               600               605
198 Tyr Asp Trp Glu Tyr Glu Gly Phe Thr Lys Cys Ser Glu Ser Cys Gly
199     610               615               620
200 Gly Gly Val Gln Glu Ala Val Val Ser Cys Leu Asn Lys Gln Thr Arg
201 625               630               635               640
202 Glu Pro Ala Glu Glu Asn Leu Cys Val Thr Ser Arg Arg Pro Pro Gln
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206 Gly Lys Trp Ser Pro Cys Ser Leu Thr Cys Gly Val Gly Leu Gln Thr
207         675              680              685
208 Arg Asp Val Phe Cys Ser His Leu Leu Ser Arg Glu Met Asn Glu Thr
209         690              695              700
210 Val Ile Leu Ala Asp Glu Leu Cys Arg Gln Pro Lys Pro Ser Thr Val
211 705              710              715              720
212 Gln Ala Cys Asn Arg Phe Asn Cys Pro Pro Ala Trp Tyr Pro Ala Gln
213             725              730              735
214 Trp Gln Pro Cys Ser Arg Thr Cys Gly Gly Gly Val Gln Lys Arg Glu
215         740              745              750
216 Val Leu Cys Lys Gln Arg Met Ala Asp Gly Ser Phe Leu Glu Leu Pro
217         755              760              765
218 Glu Thr Phe Cys Ser Ala Ser Lys Pro Ala Cys Gln Gln Ala Cys Lys
219     770              775              780
220 Lys Asp Asp Cys Pro Ser Glu Trp Leu Leu Ser Asp Trp Thr Glu Cys
221 785              790              795              800
222 Ser Thr Ser Cys Gly Glu Gly Thr Gln Thr Arg Ser Ala Ile Cys Arg
223             805              810              815
224 Lys Met Leu Lys Thr Gly Leu Ser Thr Val Val Asn Ser Thr Leu Cys
225         820              825              830
226 Pro Pro Leu Pro Phe Ser Ser Ser Ile Arg Pro Cys Met Leu Ala Thr
227         835              840              845
228 Cys Ala Arg Pro Gly Arg Pro Ser Thr Lys His Ser Pro His Ile Ala
229     850              855              860
230 Ala Ala Arg Lys Val Tyr Ile Gln Thr Arg Arg Gln Arg Lys Leu His
231 865              870              875              880
232 Phe Val Val Gly Gly Phe Ala Tyr Leu Leu Pro Lys Thr Ala Val Val
233             885              890              895
234 Leu Arg Cys Pro Ala Arg Arg Val Arg Lys Pro Leu Ile Thr Trp Glu
235         900              905              910
236 Lys Asp Gly Gln His Leu Ile Ser Ser Thr His Val Thr Val Ala Pro
237         915              920              925
238 Phe Gly Tyr Leu Lys Ile His Arg Leu Lys Pro Ser Asp Ala Gly Val
239     930              935              940
240 Tyr Thr Cys Ser Ala Gly Pro Ala Arg Glu His Phe Val Ile Lys Leu
241 945              950              955              960
242 Ile Gly Gly Asn Arg Lys Leu Val Ala Arg Pro Leu Ser Pro Arg Ser
243             965              970              975
244 Glu Glu Glu Val Leu Ala Gly Arg Lys Gly Gly Pro Lys Glu Ala Leu
245         980              985              990
246 Gln Thr His Lys His Gln Asn Gly Ile Phe Ser Asn Gly Ser Lys Ala
247         995              1000             1005
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L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date